

SEQUENCE LISTING



110> Northwest Biotherapeutics, Inc.

<120> METHODS FOR THE DIAGNOSIS AND TREATMENT OF METASTATIC PROSTATE TUMORS

<130> 8511-025-228

<140> PCT/US99/08079

<141> 1999-04-13

<160> 16

<170> PatentIn Ver. 2.0

<210> 1

<211> 4450

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (22) .. (3915)

<400> 1

acccacgcgc agcggccgga g atg cag cgg ggc gcc gcg ctg tgc ctg cga 5:

Met Gln Arg Gly Ala Ala Leu Cys Leu Arg

1 5 10

ctg tgg ctc tgc ctg gga ctc ctg gac ggc ctg gtg agt gac tac tcc 99
Leu Trp Leu Cys Leu Gly Leu Leu Asp Gly Leu Val Ser Asp Tyr Ser
15 20 25

atg acc ccc ccg acc ttg aac atc acg gag gag tca cac gtc atc gac

Met Thr Pro Pro Thr Leu Asn Ile Thr Glu Glu Ser His Val Ile Asp

30

35

40

acc ggt gac agc ctg tcc atc tcc tgc agg gga cag cac ccc ctc gag 195
Thr Gly Asp Ser Leu Ser Ile Ser Cys Arg Gly Gln His Pro Leu Glu
45 50 55

tgg gct tgg cca gga gct cag gag gcg cca gcc acc gga gac aag gac 243

Trp Ala Trp Pro Gly Ala Gln Glu Ala Pro Ala Thr Gly Asp Lys Asp

60

65

70

agc gag gac acg ggg gtg gtg cga gac tgc gag ggc aca gac gcc agg 291 Ser Glu Asp Thr Gly Val Val Arg Asp Cys Glu Gly Thr Asp Ala Arg

ccc tac tgc aag gtg ttg ctg ctg cac gag gta cat gcc aac gac aca

Pro Tyr Cys Lys Val Leu Leu Leu His Glu Val His Ala Asn Asp Thr

95

100

105

ggc agc tac gtc tgc tac tac aag tac atc aag gca cgc atc gag ggc 387
Gly Ser Tyr Val Cys Tyr Tyr Lys Tyr Ile Lys Ala Arg Ile Glu Gly
110 115 120

	g gcc ir Ala 125														435
	c aac le Asn 10														483
	g ccc al Pro														531
_	aa agc ln Ser	_		_								_			579
-	gg cgg cg Arg		_				_		_	-					627
	g cag eu Gln 205														675
	c ctg ne Leu 20														723
	cc agg co Arg														771
	gc acc /s Thr														819
gac ta Asp Ty			_	_	_				aag	tgg	gtg				867
		270	•		1114	Giu	Arg 275	Gly	Lys	Trp	Val	280	Giu	Arg	
cgc to	cc caa er Gln 285	cag	acc	cac	aca	gaa	275 ctc	tcc	agc	atc	ctg	280 acc	atc	cac	915
Arg Se	er Gln 285 cc agc al Ser	cag Gln cag	acc Thr	cac His	aca Thr	gaa Glu 290 ggc	275 ctc Leu tcg	tcc Ser	agc Ser gtg	atc Ile tgc	ctg Leu 295 aag	acc Thr	atc Ile aac	cac His	915 963
aac gt Asn Va 30	er Gln 285 cc agc al Ser	cag Gln cag Gln	acc Thr cac His	cac His gac Asp	aca Thr ctg Leu 305	gaa Glu 290 ggc Gly	ctc Leu tcg Ser	tcc Ser tat Tyr	agc Ser gtg Val	atc Ile tgc Cys 310	ctg Leu 295 aag Lys	acc Thr gcc Ala	atc Ile aac Asn	cac His aac Asn	

	a gga a Gly															1107
	g ccc o Pro			_			_	_		_	_	_			_	1155
	c agt s Ser 380			_	_			_					_	-		1203
	c acc y Thr 5				_	_				_	-		_		_	1251
	c atc n Ile															1299
	g gag s Glu															1347
	c acc u Thr															1395
	c tgg s Trp 460															1443
_	g cgg g Arg 5		_	_		_		_		_	_		_			1491
-	g gtg a Val		_	-	_	_										1539
	g acc p Thr															1587
	c cag e Gln															1635
	g gtg s Val 540		_	_												1683
	c gac o Asp 5															1731
gg	c cag	ccg	gtg	ctc	ctg	agc	tgc	caa	gcc	gac	agc	tac	aag	tac	gag	1779

.

•

Gly	Gln	Pro	Val	Leu 575	Leu	Ser	Cys	Gln	Ala 580	Asp	Ser	Tyr	Lys	Tyr 585	Glu	
						ctc Leu										1827
						gac Asp										1875
						gag Glu 625										1923
						ccc Pro										1971
						gac Asp										2019
						cag Gln										2067
						gtg Val										2115
						cac His 705										2163
						aag Lys										2211
						cgc Arg										2259
						ccc Pro										2307
						gag Glu					Met					2355
						atc Ile 785										2403
						agg Arg										2451

.

														•						
795					800					805					810					
																2499			,	
																2547				
																2595				
																2643				
																2691				
_		_	_	_												2739				
			_						_		_	_				2787				
																2835				
																2883				
																2931				
																2979				
												Gln	Glu			3027				
	Leu	Trp				Leu	Thr				Leu	Val				3075				
Phe	Gln				Gly	Met				Ala	Ser					3123				
	ggc gly caa glu galu gtl actra ctcu ctcu ctcu ctcu ggls cgg acy ctcu ctcu galus carg ttce	ggc tac Gly Tyr caa tgc Gln Cys gag cgg Glu Arg Yal 860 acc gtg Thr Val 875 cgc gcg Arg Ala ctc aac Leu Asn ctc atg Leu Met ctg cgc Arg Arg Arg Arg Arg ctg Arg Arg ttc cag ttc cag	ggc tac ctg Gly Tyr Leu Caa tgc gaa Gln Cys Glu gag cgg ctg Glu 845 gtg gtg gaa Val 860 acc gtg gcc Thr Val Ala 875 cgc gcg ctg Arg Ala Leu ctc aac gtg Leu Asn Val ctc atg gtg Leu Asn Val gag cag gcc Leu Arg Ala 940 gag agg cgc Glu Arg	ggc tac ctg tec Ser Caa tgc Glu Cys Glu Cae His 845 Clu Cae Chi Rang Chan 860 Chan 8	ggc tac ctg tec atc alc alc alc alc alc alc alc alc alc al	ggc tac ctg ser lie lie lie ggg ctg cas ggg ggg ggg ggg ggg ggg ggg ggg ggg g	ggc tac ctg tcc lile lile Met Ser lile lile lile lile lile lile lile li	ggc tac ctg ser lie lie Met Asp sac glu cys glu lie lie Met Asp sac glu cys glu lie	ggc Glytac Tyrctg Leuscc Ine Saratc Ine <br< td=""><td>ggc tac tac atc atc atc atc atc ggg g</td><td>ggc tac tac tac atc atc atc atc gac cac gly gly<td>Gly Tay Leu Sec Ite atc atc<td>ggc tac tac graph tac graph at all graph at all graph call graph ggg gag gag gag gag gag gag gag gag gag</td><td>ggc tac ctg tcc gaa atc clg atc clg</td><td>ggc tac ctg tcc atc atc atc atc atc atc atc atc atc atc</td><td>ggc tac ctg tcc atc atc atc atc atc atc atc atc atc atc</td><td>gg tac ctg tcc atc atc atc atg gac ccg gg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gau gag gag gag gag gtg cct ctg gau gag gag gag gag gag gag gag gag ga</td><td>ggc tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag 2499 Gly Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu Glu Glu Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu Glu Glu Cys Glu Tyr Leu Ser Tyr Asp Ala Ser Gln Trp Glu Phe Pro Arg 830 gag cgg ctg cac ctg ggg aag gtg ctc ggc tac ggc gcc ttc ggg aag Clu Arg Leu Gly Arg Val Leu Gly Tyr Gly Ala Phe Gly Lys 855 gtg gtg gaa gcc tcc gct ttc ggc atc cac aag ggc agc agc tgt gac Val Val Glu Ala Ser Ala Phe Gly Ile His Lys Gly Ser Ser Cys Asp 860 acc gtg gcc gtg aaa atg ctg aaa agg ggc gcc acg agc agc agc tgt gac Arg Ala Leu Met Ser Glu Leu Lys Glu Gly Ala Thr Ala Ser Glu Gln 875 ccc gcg ctg atg tcg gag ctc aag atc ctc aca atc ggc aac cac Arg Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly Aen His 895 ctc aac gtg gtc aac ctc ctc ggg gg gg tg acc aag ccc aag gcc acc Arg Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly Aen His 895 ctc aac gtg gtg atc gtg gag ttc tgc aag tac cy aac cac ag gcc ag gg ccc ctc atg gtg atc gtg gag ttc tgc aag tac cy and cy and his 895 ctc atg gtg at cac ctc ctc ggg gg gct gc acc aac ccc acc ctc atg gtg atc gtg gag ttc tgc aag tac cy 896 ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc ctc att gtg atc gtg gag ttc tgc aag tac ggc tgc acc aag ccc cac ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc ctc atg gtg atc gtg gag ttc tgc aag tac ggc tgc acc aag ccc cac ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc acl gtg gac acc gcc gag acc gcc ttc agc ccc tgc gcg gag aag tct ccc ctc atg gtg atc gtg gag tcc gc ctc agc ccc tgc gcg gag acc tcc ctc atg gtg atc gtg acc gcc tcc agc ccc atg gg ctc gcc acc ctc atg gtg atc gra gag acc acc tcc ccc gcc atg gtg gac tcc gcc agg ctg tacc ctg cgc gcc aag cgg gac acc gcc acc acc ac</td><td>ggc tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gag gag gag gag gag gag gag gag gag g</td><td>gge tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag 2499 Caa tgc gaa tac ctg tcc tac gat gcc agc cag tgg gaa ttc ccc cga all clu Glu Res Res Res Res Res Res Res Res Res Res</td></td></td></br<>	ggc tac tac atc atc atc atc atc ggg g	ggc tac tac tac atc atc atc atc gac cac gly gly <td>Gly Tay Leu Sec Ite atc atc<td>ggc tac tac graph tac graph at all graph at all graph call graph ggg gag gag gag gag gag gag gag gag gag</td><td>ggc tac ctg tcc gaa atc clg atc clg</td><td>ggc tac ctg tcc atc atc atc atc atc atc atc atc atc atc</td><td>ggc tac ctg tcc atc atc atc atc atc atc atc atc atc atc</td><td>gg tac ctg tcc atc atc atc atg gac ccg gg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gau gag gag gag gag gtg cct ctg gau gag gag gag gag gag gag gag gag ga</td><td>ggc tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag 2499 Gly Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu Glu Glu Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu Glu Glu Cys Glu Tyr Leu Ser Tyr Asp Ala Ser Gln Trp Glu Phe Pro Arg 830 gag cgg ctg cac ctg ggg aag gtg ctc ggc tac ggc gcc ttc ggg aag Clu Arg Leu Gly Arg Val Leu Gly Tyr Gly Ala Phe Gly Lys 855 gtg gtg gaa gcc tcc gct ttc ggc atc cac aag ggc agc agc tgt gac Val Val Glu Ala Ser Ala Phe Gly Ile His Lys Gly Ser Ser Cys Asp 860 acc gtg gcc gtg aaa atg ctg aaa agg ggc gcc acg agc agc agc tgt gac Arg Ala Leu Met Ser Glu Leu Lys Glu Gly Ala Thr Ala Ser Glu Gln 875 ccc gcg ctg atg tcg gag ctc aag atc ctc aca atc ggc aac cac Arg Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly Aen His 895 ctc aac gtg gtc aac ctc ctc ggg gg gg tg acc aag ccc aag gcc acc Arg Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly Aen His 895 ctc aac gtg gtg atc gtg gag ttc tgc aag tac cy aac cac ag gcc ag gg ccc ctc atg gtg atc gtg gag ttc tgc aag tac cy and cy and his 895 ctc atg gtg at cac ctc ctc ggg gg gct gc acc aac ccc acc ctc atg gtg atc gtg gag ttc tgc aag tac cy 896 ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc ctc att gtg atc gtg gag ttc tgc aag tac ggc tgc acc aag ccc cac ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc ctc atg gtg atc gtg gag ttc tgc aag tac ggc tgc acc aag ccc cac ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc acl gtg gac acc gcc gag acc gcc ttc agc ccc tgc gcg gag aag tct ccc ctc atg gtg atc gtg gag tcc gc ctc agc ccc tgc gcg gag acc tcc ctc atg gtg atc gtg acc gcc tcc agc ccc atg gg ctc gcc acc ctc atg gtg atc gra gag acc acc tcc ccc gcc atg gtg gac tcc gcc agg ctg tacc ctg cgc gcc aag cgg gac acc gcc acc acc ac</td><td>ggc tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gag gag gag gag gag gag gag gag gag g</td><td>gge tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag 2499 Caa tgc gaa tac ctg tcc tac gat gcc agc cag tgg gaa ttc ccc cga all clu Glu Res Res Res Res Res Res Res Res Res Res</td></td>	Gly Tay Leu Sec Ite atc atc <td>ggc tac tac graph tac graph at all graph at all graph call graph ggg gag gag gag gag gag gag gag gag gag</td> <td>ggc tac ctg tcc gaa atc clg atc clg</td> <td>ggc tac ctg tcc atc atc atc atc atc atc atc atc atc atc</td> <td>ggc tac ctg tcc atc atc atc atc atc atc atc atc atc atc</td> <td>gg tac ctg tcc atc atc atc atg gac ccg gg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gau gag gag gag gag gtg cct ctg gau gag gag gag gag gag gag gag gag ga</td> <td>ggc tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag 2499 Gly Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu Glu Glu Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu Glu Glu Cys Glu Tyr Leu Ser Tyr Asp Ala Ser Gln Trp Glu Phe Pro Arg 830 gag cgg ctg cac ctg ggg aag gtg ctc ggc tac ggc gcc ttc ggg aag Clu Arg Leu Gly Arg Val Leu Gly Tyr Gly Ala Phe Gly Lys 855 gtg gtg gaa gcc tcc gct ttc ggc atc cac aag ggc agc agc tgt gac Val Val Glu Ala Ser Ala Phe Gly Ile His Lys Gly Ser Ser Cys Asp 860 acc gtg gcc gtg aaa atg ctg aaa agg ggc gcc acg agc agc agc tgt gac Arg Ala Leu Met Ser Glu Leu Lys Glu Gly Ala Thr Ala Ser Glu Gln 875 ccc gcg ctg atg tcg gag ctc aag atc ctc aca atc ggc aac cac Arg Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly Aen His 895 ctc aac gtg gtc aac ctc ctc ggg gg gg tg acc aag ccc aag gcc acc Arg Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly Aen His 895 ctc aac gtg gtg atc gtg gag ttc tgc aag tac cy aac cac ag gcc ag gg ccc ctc atg gtg atc gtg gag ttc tgc aag tac cy and cy and his 895 ctc atg gtg at cac ctc ctc ggg gg gct gc acc aac ccc acc ctc atg gtg atc gtg gag ttc tgc aag tac cy 896 ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc ctc att gtg atc gtg gag ttc tgc aag tac ggc tgc acc aag ccc cac ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc ctc atg gtg atc gtg gag ttc tgc aag tac ggc tgc acc aag ccc cac ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc acl gtg gac acc gcc gag acc gcc ttc agc ccc tgc gcg gag aag tct ccc ctc atg gtg atc gtg gag tcc gc ctc agc ccc tgc gcg gag acc tcc ctc atg gtg atc gtg acc gcc tcc agc ccc atg gg ctc gcc acc ctc atg gtg atc gra gag acc acc tcc ccc gcc atg gtg gac tcc gcc agg ctg tacc ctg cgc gcc aag cgg gac acc gcc acc acc ac</td> <td>ggc tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gag gag gag gag gag gag gag gag gag g</td> <td>gge tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag 2499 Caa tgc gaa tac ctg tcc tac gat gcc agc cag tgg gaa ttc ccc cga all clu Glu Res Res Res Res Res Res Res Res Res Res</td>	ggc tac tac graph tac graph at all graph at all graph call graph ggg gag gag gag gag gag gag gag gag gag	ggc tac ctg tcc gaa atc clg atc clg	ggc tac ctg tcc atc	ggc tac ctg tcc atc	gg tac ctg tcc atc atc atc atg gac ccg gg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gau gag gag gag gag gtg cct ctg gau gag gag gag gag gag gag gag gag ga	ggc tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag 2499 Gly Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu Glu Glu Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu Glu Glu Cys Glu Tyr Leu Ser Tyr Asp Ala Ser Gln Trp Glu Phe Pro Arg 830 gag cgg ctg cac ctg ggg aag gtg ctc ggc tac ggc gcc ttc ggg aag Clu Arg Leu Gly Arg Val Leu Gly Tyr Gly Ala Phe Gly Lys 855 gtg gtg gaa gcc tcc gct ttc ggc atc cac aag ggc agc agc tgt gac Val Val Glu Ala Ser Ala Phe Gly Ile His Lys Gly Ser Ser Cys Asp 860 acc gtg gcc gtg aaa atg ctg aaa agg ggc gcc acg agc agc agc tgt gac Arg Ala Leu Met Ser Glu Leu Lys Glu Gly Ala Thr Ala Ser Glu Gln 875 ccc gcg ctg atg tcg gag ctc aag atc ctc aca atc ggc aac cac Arg Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly Aen His 895 ctc aac gtg gtc aac ctc ctc ggg gg gg tg acc aag ccc aag gcc acc Arg Ala Leu Met Ser Glu Leu Lys Ile Leu Ile His Ile Gly Aen His 895 ctc aac gtg gtg atc gtg gag ttc tgc aag tac cy aac cac ag gcc ag gg ccc ctc atg gtg atc gtg gag ttc tgc aag tac cy and cy and his 895 ctc atg gtg at cac ctc ctc ggg gg gct gc acc aac ccc acc ctc atg gtg atc gtg gag ttc tgc aag tac cy 896 ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc ctc att gtg atc gtg gag ttc tgc aag tac ggc tgc acc aag ccc cac ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc ctc atg gtg atc gtg gag ttc tgc aag tac ggc tgc acc aag ccc cac ctc atg gtg atc gtg gag ttc tgc aag tac ggc acc ctc tcc aac ttc acl gtg gac acc gcc gag acc gcc ttc agc ccc tgc gcg gag aag tct ccc ctc atg gtg atc gtg gag tcc gc ctc agc ccc tgc gcg gag acc tcc ctc atg gtg atc gtg acc gcc tcc agc ccc atg gg ctc gcc acc ctc atg gtg atc gra gag acc acc tcc ccc gcc atg gtg gac tcc gcc agg ctg tacc ctg cgc gcc aag cgg gac acc gcc acc acc ac	ggc tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gag gag gag gag gtg cct ctg gag gag gag gag gag gag gag gag gag g	gge tac ctg tcc atc atc atg gac ccc ggg gag gtg cct ctg gag gag gag gtg cct ctg gag gag gag 2499 Caa tgc gaa tac ctg tcc tac gat gcc agc cag tgg gaa ttc ccc cga all clu Glu Res

His Arg Asp Leu Ala		ctg ctg tcg gaa agc gac gtg 317: Leu Leu Ser Glu Ser Asp Val 1045 1050	1
		cgg gac atc tac aaa gac ccc 3219 Arg Asp Ile Tyr Lys Asp Pro 1060 1065	9
	••	ctg ccc ctg aag tgg atg gcc 326° Leu Pro Leu Lys Trp Met Ala 1080	7
		acc acg cag agt gac gtg tgg 3319 Thr Thr Gln Ser Asp Val Trp 1095	5
		ttc tct ctg ggg gcc tcc ccg 3365 Phe Ser Leu Gly Ala Ser Pro 1110	3
Tyr Pro Gly Val Gln		ttc tgc cag cgc gtg aga gac 3411 Phe Cys Gln Arg Val Arg Asp 1125 1130	1
		gcc act ccc gcc ata cgc cac 3459 Ala Thr Pro Ala Ile Arg His 1140 1145	9
		ccc aag gcg aga cct gca ttc 350° Pro Lys Ala Arg Pro Ala Phe 1160	7
		ctg ctc cag ggc agg ggc ctg 3559 Leu Leu Gln Gly Arg Gly Leu 1175	5
		ccg cgc agc tct cag agc tca 3600 Pro Arg Ser Ser Gln Ser Ser 1190	3
Glu Glu Gly Ser Phe		acc atg gcc cta cac atc gcc 365: Thr Met Ala Leu His Ile Ala 1205 1210	1
	Asp Ser Pro Pro	agc ctg cag cgc cac agc ctg 3699 Ser Leu Gln Arg His Ser Leu 1220 1225	9
	-	ttt ccc ggg tgc ctg gcc aga 374° Phe Pro Gly Cys Leu Ala Arg 1240	7
		atg aag aca ttt gag gaa ttc 3799 Met Lys Thr Phe Glu Glu Phe	5

ccc atg acc cca acg acc tac aaa ggc tct gtg gac aac cag aca gac 3843 Pro Met Thr Pro Thr Thr Tyr Lys Gly Ser Val Asp Asn Gln Thr Asp 1260 1265 1270

agt ggg atg gtg ctg gcc tcg gag gag ttt gag cag ata gag agc agg 3891 Ser Gly Met Val Leu Ala Ser Glu Glu Phe Glu Gln Ile Glu Ser Arg 1275 1280 1285 1290

cat aga caa gaa agc ggc ttc agg tagctgaagc agagagagag aaggcagcat 3945 His Arg Gln Glu Ser Gly Phe Arg 1295

acgtcagcat tttcttctct gcacttataa gaaagatcaa agactttaag actttcgcta 4005
tttcttctac tgctatctac tacaaacttc aaagaggaac caggaggaca agaggagcat 4065
gaaagtggac aaggagtgtg accactgaag caccacaggg aggggttagg cctccggatg 4125
actgcgggca ggcctggata atatccagcc tcccacaaga agctggtgga gcagagtgtt 4185
ccctgactcc tccaaggaaa gggagacgcc ctttcatggt ctgctgagta acaggtgcct 4245
tcccagacac tggcgttact gcttgaccaa agagccctca agcggccctt atgccagcgt 4305
gacagagggc tcacctcttg ccttctaggt cacttctcac aatgtccctt cagcacctga 4365
ccctgtgccc gccgattatt ccttggtaat atgagtaata catcaaagag tagtattaaa 4425
agctaattaa tcatgtttat aaaaa

<210> 2

<211> 1298

<212> PRT

<213> Homo sapiens

<400> 2

Met Gln Arg Gly Ala Ala Leu Cys Leu Arg Leu Trp Leu Cys Leu Gly
1 5 10 15

Leu Leu Asp Gly Leu Val Ser Asp Tyr Ser Met Thr Pro Pro Thr Leu 20 25 30

Asn Ile Thr Glu Glu Ser His Val Ile Asp Thr Gly Asp Ser Leu Ser

Ile Ser Cys Arg Gly Gln His Pro Leu Glu Trp Ala Trp Pro Gly Ala
50 55 60

Gln Glu Ala Pro Ala Thr Gly Asp Lys Asp Ser Glu Asp Thr Gly Val
65 70 75 80

Val Arg Asp Cys Glu Gly Thr Asp Ala Arg Pro Tyr Cys Lys Val Leu 85 90 95

Leu Leu His Glu Val His Ala Asn Asp Thr Gly Ser Tyr Val Cys Tyr
100 105 110

Tyr	Lys	Tyr 115	Ile	Lys	Ala	Arg	Ile 120	Glu	Gly	Thr	Thr	Ala 125	Ala	Ser	Ser
Tyr	Val 130	Phe	Val	Arg	Asp	Phe 135	Glu	Gln	Pro	Phe	Ile 140	Asn	Lys	Pro	Asp
Thr 145	Leu	Leu	Val	Asn	Arg 150	Lys	Asp	Ala	Met	Trp 155	Val	Pro	Cys	Leu	Val 160
Ser	Ile	Pro	Gly	Leu 165	Asn	Val	Thr	Leu	Arg 170	Ser	Gln	Ser	Ser	Val 175	Leu
Trp	Pro	Asp	Gly 180	Gln	Glu	Val	Val	Trp 185	Asp	Asp	Arg	Arg	Gly 190	Met	Leu
Val	Ser	Thr 195	Pro	Leu	Leu	His	Asp 200	Ala	Leu	Tyr	Leu	Gln 205	Cys	Glu	Thr
Thr	Trp 210	Gly	Asp	Gln	Asp	Phe 215	Leu	Ser	Asn	Pro	Phe 220	Leu	Val	His	Ile
225	_				230	_				235				Ser	240
				245				•	250					Trp 255	-
			260					265					270	Lys	
		275					280					285		Thr	
	290					295					300			His	- ,
305					310					315				Phe	320
				325					330					Val 335	•
			340					345					350	Leu	
		355		-			360					365		Gln	
_	370					375			_		380			Ala	
385					390					395				Leu	400
Leu	Trp	Asn	ser	Ala 405	Ala	GTÀ	ьeu	Arg	Arg 410	Asn	шe	ser	Leu	Glu 415	ьеu
															•

. .

- Val Val Asn Val Pro Pro Gln Ile His Glu Lys Glu Ala Ser Ser Pro 420 425 430
- Ser Ile Tyr Ser Arg His Ser Arg Gln Ala Leu Thr Cys Thr Ala Tyr 435 440 445
- Gly Val Pro Leu Pro Leu Ser Ile Gln Trp His Trp Arg Pro Trp Thr 450 455 460
- Pro Cys Lys Met Phe Ala Gln Arg Ser Leu Arg Arg Arg Gln Gln Gln 465 470 475 480
- Asp Leu Met Pro Gln Cys Arg Asp Trp Arg Ala Val Thr Thr Gln Asp 485 490 495
- Ala Val Asn Pro Ile Glu Ser Leu Asp Thr Trp Thr Glu Phe Val Glu 500 505 510
- Gly Lys Asn Lys Thr Val Ser Lys Leu Val Ile Gln Asn Ala Asn Val 515 520 525
- Ser Ala Met Tyr Lys Cys Val Val Ser Asn Lys Val Gly Gln Asp Glu 530 540
- Arg Leu Ile Tyr Phe Tyr Val Thr Thr Ile Pro Asp Gly Phe Thr Ile 545 550 555 560
- Glu Ser Lys Pro Ser Glu Glu Leu Leu Glu Gly Gln Pro Val Leu Leu 565 570 575
- Ser Cys Gln Ala Asp Ser Tyr Lys Tyr Glu His Leu Arg Trp Tyr Arg 580 585 590
- Leu Asn Leu Ser Thr Leu His Asp Ala His Gly Asn Pro Leu Leu 595 600 605
- Asp Cys Lys Asn Val His Leu Phe Ala Thr Pro Leu Ala Ala Ser Leu 610 620
- Glu Glu Val Ala Pro Gly Ala Arg His Ala Thr Leu Ser Leu Ser Ile 625 630 635 640
- Pro Arg Val Ala Pro Glu His Glu Gly His Tyr Val Cys Glu Val Gln
 645 650 655
- Asp Arg Arg Ser His Asp Lys His Cys His Lys Lys Tyr Leu Ser Val 660 665 670
- Gln Ala Leu Glu Ala Pro Arg Leu Thr Gln Asn Leu Thr Asp Leu Leu 675 680 685
- Val Asn Val Ser Asp Ser Leu Glu Met Gln Cys Leu Val Ala Gly Ala 690 695 700
- His Ala Pro Ser Ile Val Trp Tyr Lys Asp Glu Arg Leu Leu Glu Glu 705 710 715 720

- Lys Ser Gly Val Asp Leu Ala Asp Ser Asn Gln Lys Leu Ser Ile Gln 725 730 735
- Arg Val Arg Glu Glu Asp Ala Gly Pro Tyr Leu Cys Ser Val Cys Arg
 740 745 750
- Pro Lys Gly Cys Val Asn Ser Ser Ala Ser Val Ala Val Glu Gly Ser
 755 760 765
- Glu Asp Lys Gly Ser Met Glu Ile Val Ile Leu Val Gly Thr Gly Val
 770 780
- Ile Ala Val Phe Phe Trp Val Leu Leu Leu Leu Ile Phe Cys Asn Met 785 790 795 800
- Arg Arg Pro Ala His Ala Asp Ile Lys Thr Gly Tyr Leu Ser Ile Ile 805 810 815
- Met Asp Pro Gly Glu Val Pro Leu Glu Glu Gln Cys Glu Tyr Leu Ser 820 825 830
- Tyr Asp Ala Ser Gln Trp Glu Phe Pro Arg Glu Arg Leu His Leu Gly 835 840 845
- Arg Val Leu Gly Tyr Gly Ala Phe Gly Lys Val Val Glu Ala Ser Ala 850 855 860
- Phe Gly Ile His Lys Gly Ser Ser Cys Asp Thr Val Ala Val Lys Met 865 870 875 880
- Leu Lys Glu Gly Ala Thr Ala Ser Glu Gln Arg Ala Leu Met Ser Glu 885 890 895
- Leu Lys Ile Leu Ile His Ile Gly Asn His Leu Asn Val Val Asn Leu 900 905 910
- Leu Gly Ala Cys Thr Lys Pro Gln Gly Pro Leu Met Val Ile Val Glu 915 920 925
- Phe Cys Lys Tyr Gly Asn Leu Ser Asn Phe Leu Arg Ala Lys Arg Asp 930 935 940
- Ala Phe Ser Pro Cys Ala Glu Lys Ser Pro Glu Gln Arg Gly Arg Phe 945 950 955 960
- Arg Ala Met Val Glu Leu Ala Arg Leu Asp Arg Arg Pro Gly Ser 965 970 975
- Ser Asp Arg Val Leu Phe Ala Arg Phe Ser Lys Thr Glu Gly Gly Ala 980 985 990
- Arg Arg Ala Ser Pro Asp Gln Glu Ala Glu Asp Leu Trp Leu Ser Pro
 995 1000 1005
- Leu Thr Met Glu Asp Leu Val Cys Tyr Ser Phe Gln Val Ala Arg Gly
 1010 1015 1020

Met Glu Phe Leu Ala Ser Arg Lys Cys Ile His Arg Asp Leu Ala Ala 025 1030 1035 1040

Arg Asn Ile Leu Leu Ser Glu Ser Asp Val Val Lys Ile Cys Asp Phe 1045 1050 1055

Gly Leu Ala Arg Asp Ile Tyr Lys Asp Pro Asp Tyr Val Arg Lys Gly
1060 1065 1070

Ser Ala Arg Leu Pro Leu Lys Trp Met Ala Pro Glu Ser Ile Phe Asp 1075 1080 1085

Lys Val Tyr Thr Thr Gln Ser Asp Val Trp Ser Phe Gly Val Leu Leu 1090 1095 1100

Trp Glu Ile Phe Ser Leu Gly Ala Ser Pro Tyr Pro Gly Val Gln Ile 105 1110 1115 1120

Asn Glu Glu Phe Cys Gln Arg Val Arg Asp Gly Thr Arg Met Arg Ala 1125 1130 1135

Pro Glu Leu Ala Thr Pro Ala Ile Arg His Ile Met Leu Asn Cys Trp 1140 1145 1150

Ser Gly Asp Pro Lys Ala Arg Pro Ala Phe Ser Asp Leu Val Glu Ile 1155 1160 1165

Leu Gly Asp Leu Leu Gln Gly Arg Gly Leu Gln Glu Glu Glu Val 1170 1175 1180

Cys Met Ala Pro Arg Ser Ser Gln Ser Ser Glu Glu Gly Ser Phe Ser 185 1190 1195 1200

Gln Val Ser Thr Met Ala Leu His Ile Ala Gln Ala Asp Ala Glu Asp 1205 1210 1215

Ser Pro Pro Ser Leu Gln Arg His Ser Leu Ala Ala Arg Tyr Tyr Asn 1220 1225 1230

Trp Val Ser Phe Pro Gly Cys Leu Ala Arg Gly Ala Glu Thr Arg Gly 1235 1240 1245

Ser Ser Arg Met Lys Thr Phe Glu Glu Phe Pro Met Thr Pro Thr Thr 1250 1255 1260

Tyr Lys Gly Ser Val Asp Asn Gln Thr Asp Ser Gly Met Val Leu Ala 265 1270 1275 1280

Ser Glu Glu Phe Glu Gln Ile Glu Ser Arg His Arg Gln Glu Ser Gly 1285 1290 1295

Phe Arg

```
<211> 15
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:
      oligonucleotide
<400> 3
ggcgccccgc tgcat
                                                                    15
<210> 4
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<400> 4
                                                                    23
cgaagtggtg aagttcatgg atg
<210> 5
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
ttctgtatca gtctttcctg gtgag
                                                                    25
<210> 6
<211> 24
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: primer
<400> 6
                                                                    24
ctggcatggt cttctgtgaa agca
<210> 7
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<400> 7
```

```
aataccagtg gatgtgatgc gg
                                                                    22
<210> 8
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: primer
<400> 8
                                                                    20
taccacagtg tcaggcagcg
<210> 9
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<400> 9
                                                                    20
atcaaattct cggttggccc
<210> 10
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: primer
<400> 10
                                                                    20
agagggatgg agttcctggc
<210> 11
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<400> 11
                                                                    22
aataccagtg gatgtgatgc gg
<210> 12
<211> 21
<212> DNA
<213> Artificial Sequence
```

<220>

	Description of Artificial oligonucleotide	Sequence:	
<400>	12		
	gcaga aagttcatgg t		21
<210>			
<211><212>			
	Artificial Sequence		
<220>			
<223>	Description of Artificial oligonucleotide	Sequence:	
<400>	13		
caagt	gcatg gtgga		15
<210><211>			
<211>			
	Artificial Sequence		
<220>			
<223>	Description of Artificial oligonucleotide	Sequence:	
<400>	14		
cacct	gctc tgcat		15
cacct	egete tgeat		15
		,	15
<210><211>	15		15
<210>	15 14		15
<210><211><212>	15 14		15
<210><211><211><212><213>	15 14 DNA Artificial Sequence		15
<210><211><211><212><213>	15 14 DNA	Sequence:	15
<210><211><211><212><213>	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide	Sequence:	15
<210> <211> <212> <213> <223> <400>	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide	Sequence:	15
<210> <211> <212> <213> <223> <400>	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide 15	Sequence:	
<210><211><211><212><213><220><223> 400	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide 15 egtcc caga	Sequence:	
<210><211><211><212><213><220><223> 400 221	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide 15 egtcc caga	Sequence:	
<210><211><211><212><213> 223 223 400 223 400 210 211 212	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide 15 egtcc caga 16 17 DNA	Sequence:	
<210><211><211><212><213> 223 223 400 223 400 210 211 212	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide 15 egtcc caga	Sequence:	
<210><211><211><212><213> 223 223 400 223 400 210 211 212	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide 15 egtcc caga 16 17 DNA	Sequence:	
<210><211><211><212><213> 223 223 400 221 210 211 212 213 220	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide 15 egtcc caga 16 17 DNA		
<210><211><211><212><213> 223 223 400 221 210 211 212 213 220	15 14 DNA Artificial Sequence Description of Artificial oligonucleotide 15 cgtcc caga 16 17 DNA Artificial Sequence Description of Artificial oligonucleotide		

.